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<110> EXELIXIS, INC.

<120> PRKCS AS MODIFIERS OF THE BETA CATENIN PATHWAY AND METHODS OF USE

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<150> US 60/495,172

<151> 2003-08-14

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 <212> PRT
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<400> 11

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Glu Pro Ser Ile Ser Phe Glu Gly Leu Cys Asn Glu Val Arg Asp Met
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Cys Ser Phe Asp Asn Glu Gln Leu Phe Thr Met Lys Trp Ile Asp Glu
50 55 60

Glu Gly Asp Pro Cys Thr Val Ser Ser Gln Leu Glu Leu Glu Glu Ala
65 70 75 80

Phe Arg Leu Tyr Glu Leu Asn Lys Asp Ser Glu Leu Leu Ile His Val
85 90 95

Phe Pro Cys Val Pro Glu Arg Pro Gly Met Pro Cys Pro Gly Glu Asp
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Lys Ser Ile Tyr Arg Arg Gly Ala Arg Arg Trp Arg Lys Leu Tyr Cys
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130 135 140

Cys Ala Ile Cys Thr Asp Arg Ile Trp Gly Leu Gly Arg Gln Gly Tyr
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Lys Cys Ile Asn Cys Lys Leu Leu Val His Lys Lys Cys His Lys Leu
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Val Thr Ile Glu Cys Gly Arg His Ser Leu Pro Gln Glu Pro Val Met
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Pro Met Asp Gln Ser Ser Met His Ser Asp His Ala Gln Thr Val Ile
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Pro Tyr Asn Pro Ser Ser His Glu Ser Leu Asp Gln Val Gly Glu Glu
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Lys Glu Ala Met Asn Thr Arg Glu Ser Gly Lys Ala Ser Ser Ser Leu
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Gly Leu Gln Asp Phe Asp Leu Leu Arg Val Ile Gly Arg Gly Ser Tyr

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<400> 12

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Ala Ala Thr Thr Phe Glu Glu Leu Cys Glu Glu Val Arg Asp Met Cys
35 40 45

Arg Leu His Gln Gln His Pro Leu Thr Leu Lys Trp Val Asp Ser Glu
50 55 60

Gly Asp Pro Cys Thr Val Ser Ser Gln Met Glu Leu Glu Glu Ala Phe
65 70 75 80

Arg Leu Ala Arg Gln Cys Arg Asp Glu Gly Leu Ile Ile His Val Phe
85 90 95

Pro Ser Thr Pro Glu Gln Pro Gly Leu Pro Cys Pro Gly Glu Asp Lys
100 105 110

Ser Ile Tyr Arg Arg Gly Ala Arg Arg Trp Arg Lys Leu Tyr Arg Ala
115 120 125

Asn Gly His Leu Phe Gln Ala Lys Arg Phe Asn Arg Arg Ala Tyr Cys
130 135 140

Gly Gln Cys Ser Glu Arg Ile Trp Gly Leu Ala Arg Gln Gly Tyr Arg
145 150 155 160

Cys Ile Asn Cys Lys Leu Leu Val His Lys Arg Cys His Gly Leu Val
165 170 175

Pro Leu Thr Cys Arg Lys His Met Asp Ser Val Met Pro Ser Gln Glu
180 185 190

Pro Pro Val Asp Asp Lys Asn Glu Asp Ala Asp Leu Pro Ser Glu Glu
195 200 205

Thr Asp Gly Ile Ala Tyr Ile Ser Ser Ser Arg Lys His Asp Ser Ile
210 215 220

Lys Asp Asp Ser Glu Asp Leu Lys Pro Val Ile Asp Gly Met Asp Gly
225 230 235 240

Ile Lys Ile Ser Gln Gly Leu Gly Leu Gln Asp Phe Asp Leu Ile Arg
245 250 255

Val Ile Gly Arg Gly Ser Tyr Ala Lys Val Leu Leu Val Arg Leu Lys
260 265 270

Lys Asn Asp Gln Ile Tyr Ala Met Lys Val Val Lys Lys Glu Leu Val
275 280 285

His Asp Asp Glu Asp Ile Asp Trp Val Gln Thr Glu Lys His Val Phe
290 295 300

Glu Gln Ala Ser Ser Asn Pro Phe Leu Val Gly Leu His Ser Cys Phe

305		310		315		320
Gln Thr Thr Ser	Arg Leu Phe Leu Val	Ile Glu Tyr Val	Asn Gly Gly			
	325		330		335	
Asp Leu Met Phe	His Met Gln Arg	Gln Arg Lys Leu	Pro Glu Glu His			
	340		345		350	
Ala Arg Phe Tyr	Ala Ala Glu Ile	Cys Ile Ala Leu	Asn Phe Leu His			
	355		360		365	
Glu Arg Gly Ile	Ile Tyr Arg Asp	Leu Lys Leu Asp	Asn Val Leu Leu			
	370		375		380	
Asp Ala Asp Gly	His Ile Lys Leu	Thr Asp Tyr Gly	Met Cys Lys Glu			
	385		390		395	
Gly Leu Gly Pro	Gly Asp Thr Thr	Ser Thr Phe Cys	Gly Thr Pro Asn			
	405		410		415	
Tyr Ile Ala Pro	Glu Ile Leu Arg	Gly Glu Glu Tyr	Gly Phe Ser Val			
	420		425		430	
Asp Trp Trp Ala	Leu Gly Val Leu	Met Phe Glu Met	Met Ala Gly Arg			
	435		440		445	
Ser Pro Phe Asp	Ile Ile Thr Asp	Asn Pro Asp Met	Asn Thr Glu Asp			
	450		455		460	
Tyr Leu Phe Gln	Val Ile Leu Glu	Lys Pro Ile Arg	Ile Pro Arg Phe			
	465		470		475	
Leu Ser Val Lys	Ala Ser His Val	Leu Lys Gly Phe	Leu Asn Lys Asp			
	485		490		495	
Pro Lys Glu Arg	Leu Gly Cys Arg	Pro Gln Thr Gly	Phe Ser Asp Ile			
	500		505		510	
Lys Ser His Ala	Phe Phe Arg Ser	Ile Asp Trp Asp	Leu Leu Glu Lys			
	515		520		525	
Lys Gln Ala Leu	Pro Pro Phe Gln	Pro Gln Ile Thr	Asp Asp Tyr Gly			

530

535

540

Leu Asp Asn Phe Asp Thr Gln Phe Thr Ser Glu Pro Val Gln Leu Thr
545 550 555 560

Pro Asp Asp Glu Asp Ala Ile Lys Arg Ile Asp Gln Ser Glu Phe Glu
565 570 575

Gly Phe Glu Tyr Ile Asn Pro Leu Leu Leu Ser Thr Glu Glu Ser Val
580 585 590